AMENDMENTS

In the claims

Please cancel Claim 21, without prejudice.

Please amend Claims 1 and 3, as follows:

Claim 1 (Once Amended). A modified enzyme which comprises a catalytically active amino acid sequence derived from a cellulolytic enzyme EGI exhibiting the following properties:

(a) derived from *Humicola insolens* or *Trichoderma reseei*,

(b) approximate molecular weight of about 50 kDa,

(c) iso-electric point of 5.5, and

(d) containing 415 amino acid;

linked to an amino acid sequence comprising a cellulose binding domain;

wherein said modified enzyme comprises a linking region between said catalytically active amino acid sequence of a cellulolytic enzyme EGI and said amino acid sequence comprising a cellulose binding domain; further wherein said linking region is selected from the group consisting of: *Humicola insolens* family 45 cellulase linker, Nifa gene of *Klebsiella pneumoniae*-CiP linker, *E. coli* OmpA gene-CiP linker, E3 cellulase *Thermomonospora fusca* linker, CenA cellulase linker, nucleophilic polyethylene glycol derivative linker, carboxyl polyethylene glycol derivative linker, electrophilically activated polyethylene glycol derivative linker, sulfhydryl-selective polyethylene glycol derivative linker, hertofunctional polyethylene glycol derivative linker, biotin polyethylene glycol derivative linker, vinyl polyethylene glycol derivative linker, polyethylene glycol silane derivative linker, polyethylene glycol phospholipid derivative linker and mixtures thereof.

Claim 3 (Once Amended). A modified enzyme according to claim 1 wherein the amino acid sequence comprising a cellulose binding domain is selected from CBD family 45 from *Humicol insolens*, CBD CenC from *Cellulomonas fimi* and/or CBD Cellulozome from *Clostridium cellulovorans*.